0500/0280

OIPE

#2.

RAW SEQUENCE LISTING   DATE: 02/06/2002
PATENT APPLICATION: US/10/053,291   TIME: 09:32:54

Input Set : N:\Crf3\RULE60\10053291.raw
Output Set: N:\CRF3\02062002\J053291.raw

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 1 <110> APPLICANT: Stuhlmann, Heidi
 2        Xiong, Jing-Wei
 3        Taubman, Mark B.
 4 <120> TITLE OF INVENTION: VASCULAR ENDOTHELIAL ZINC FINGER 1 GENE
 5        AND PROTEIN AND USES THEREOF
 6 <130> FILE REFERENCE: 31200
 8 <140> CURRENT APPLICATION NUMBER: 10/053,291
 9 <141> CURRENT FILING DATE: 2002-01-17
11 <150> PRIOR APPLICATION NUMBER: US/09/083,290A
12 <151> PRIOR FILING DATE: 1998-05-22
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16 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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25        ctgcccctcc tgagttctgc tgtggagccc cctgatcaga aaccgttgct tccaatacca    180
26        attactcaga aacctcaggc tgcaccagaa acattaaagg atgccattgg gattaaaaaa    240
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42        gaagaagctg tcaaagcaag aaagaaagaa gctgccaacc tgtgccaaac ctccacggct   1200
43        gctacgacac cagtgactct cactactcca ttcaatataa cgtcctctgt gtcgtctggg   1260
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93                      20                  25                  30
94      Ser Ala Val Glu Pro Pro Asp Gln Lys Pro Leu Leu Pro Ile Pro Ile
95                  35                  40                  45
96      Thr Gln Lys Pro Gln Ala Ala Pro Glu Thr Leu Lys Asp Ala Ile Gly
97              50                  55                  60
98      Ile Lys Lys Glu Lys Pro Lys Thr Ser Phe Val Cys Thr Tyr Cys Ser
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Input Set : N:\Crf3\RULE60\10053291.raw
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103				100					105					110		
104	Val	Pro	Leu	Ile	Ser	Thr	Ile	Ala	Gly	Asp	Ser	Ser	Arg	Thr	Ser	Leu
105			115					120					125			
106	Val	Ser	Thr	Ile	Ala	Gly	Ile	Leu	Ser	Thr	Val	Thr	Thr	Ser	Ser	Ser
107			130					135				140				
108	Gly	Thr	Asn	Pro	Ser	Ser	Ser	Ala	Ser	Thr	Thr	Ala	Met	Pro	Val	Pro
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110	Gln	Ser	Val	Lys	Lys	Pro	Ser	Lys	Pro	Val	Lys	Lys	Asn	His	Ala	Cys
111				165					170					175		
112	Glu	Met	Cys	Gly	Lys	Ala	Phe	Arg	Asp	Val	Tyr	His	Leu	Asn	Arg	His
113				180					185					190		
114	Lys	Leu	Ser	His	Ser	Asp	Glu	Lys	Pro	Phe	Glu	Cys	Pro	Ile	Cys	Asn
115			195					200					205			
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117		210					215					220				
118	Glu	Gly	Gly	Ile	Thr	Lys	Pro	Tyr	Thr	Cys	Ser	Val	Cys	Gly	Lys	Gly
119		225					230					235					240
120	Phe	Ser	Arg	Pro	Asp	His	Leu	Ser	Cys	His	Val	Lys	His	Val	His	Ser
121				245					250					255		
122	Thr	Glu	Arg	Pro	Phe	Lys	Cys	Gln	Thr	Cys	Thr	Ala	Ala	Phe	Ala	Thr
123			260					265					270			
124	Lys	Asp	Arg	Leu	Arg	Thr	His	Met	Val	Arg	His	Glu	Gly	Lys	Val	Ser
125			275					280					285			
126	Cys	Asn	Ile	Cys	Gly	Lys	Leu	Leu	Ser	Ala	Ala	Tyr	Ile	Thr	Ser	His
127		290					295					300				
128	Leu	Lys	Thr	His	Gly	Gln	Ser	Gln	Ser	Ile	Asn	Cys	Asn	Thr	Cys	Lys
129		305					310					315					320
130	Gln	Gly	Ile	Ser	Lys	Thr	Cys	Met	Ser	Glu	Glu	Thr	Ser	Asn	Gln	Lys
131				325					330					335		
132	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	His
133				340					345					350		
134	Val	Thr	Ser	Trp	Pro	Gly	Lys	Gln	Val	Glu	Thr	Leu	Arg	Leu	Trp	Glu
135			355					360					365			
136	Glu	Ala	Val	Lys	Ala	Arg	Lys	Lys	Glu	Ala	Ala	Asn	Leu	Cys	Gln	Thr
137			370					375					380			
138	Ser	Thr	Ala	Ala	Thr	Thr	Pro	Val	Thr	Leu	Thr	Thr	Pro	Phe	Asn	Ile
139		385					390					395					400
140	Thr	Ser	Ser	Val	Ser	Ser	Gly	Thr	Met	Ser	Asn	Pro	Val	Thr	Val	Ala
141				405					410					415		
142	Ala	Ala	Met	Ser	Met	Arg	Ser	Pro	Val	Asn	Val	Ser	Ser	Ala	Val	Asn
143				420					425					430		
144	Ile	Thr	Ser	Pro	Leu	Ala	Met	Thr	Ser	Pro	Leu	Thr	Leu	Thr	Thr	Pro
145			435					440					445			
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196      gtgataccag ctctgatatg caaagcatat gataatttat cattctaact tcaacgtata    2220
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208                      20                  25                  30
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210                      35                  40                  45
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212          50                  55                  60
213      Ile Lys Lys Glu Lys Pro Lys Thr Ser Phe Val Cys Thr Tyr Cys Ser
214      65                  70                  75                  80
215      Lys Ala Phe Arg Asp Ser Tyr His Leu Arg Arg His Glu Ser Cys His
216                      85                  90                  95
217      Thr Gly Ile Lys Leu Val Ser Arg Pro Lys Lys Thr Pro Thr Thr Val
218                      100                 105                 110
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222          130                 135                 140
223      Gly Thr Asn Pro Ser Ser Ser Ala Ser Thr Thr Ala Met Pro Val Thr
224      145                 150                 155                 160
225      Gln Ser Val Lys Lys Pro Ser Lys Pro Val Lys Lys Asn His Ala Cys
226                      165                 170                 175
227      Glu Met Cys Gly Lys Ala Phe Arg Asp Val Tyr His Leu Asn Arg His
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229      Lys Leu Ser His Ser Asp Glu Lys Pro Phe Glu Cys Pro Ile Cys Asn
230                      195                 200                 205
231      Gln Arg Phe Lys Arg Lys Asp Arg Met Thr Tyr His Val Arg Ser His
232          210                 215                 220
233      Glu Gly Gly Ile Thr Lys Pro Tyr Thr Cys Ser Val Cys Gly Lys Gly
234      225                 230                 235                 240
235      Phe Ser Arg Pro Asp His Leu Ser Cys His Val Lys His Val His Ser
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238                      260                 265                 270
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241      Cys Asn Ile Cys Gly Lys Leu Leu Ser Ala Ala Tyr Ile Thr Ser His
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243      Leu Lys Thr His Gly Gln Ser Gln Ser Ile Asn Cys Asn Thr Cys Lys
244      305                 310                 315                 320
245      Gln Gly Ile Ser Lys Thr Cys Met Ser Glu Glu Thr Ser Asn Gln Lys
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